

Please amend claims 4, 8, and 24 as follows:

4. (Twice Amended) An isolated nucleic acid molecule consisting of a nucleotide sequence selected from the group consisting of:

- C<sup>2</sup>
- (a) a nucleotide sequence that encodes a polypeptide comprising the amino acid sequence of SEQ ID NO:2;
  - (b) a nucleotide sequence consisting of SEQ ID NO:1;
  - (c) a nucleotide sequence consisting of SEQ ID NO:3; and
  - (d) a nucleotide sequence that is completely complementary over the entire length of a nucleotide sequence of (a)-(c).

C<sup>3</sup> 8. (Twice Amended) A vector comprising the nucleic acid molecule of claim 4.

C<sup>4</sup> 24. (Amended) A process for producing a polypeptide comprising the amino acid sequence of SEQ ID NO:2, the process comprising culturing the host cell of claim 9 under conditions sufficient for the production of said polypeptide, and recovering said polypeptide, thereby producing said polypeptide.

**In the Specification:**

Please replace the last two consecutive paragraphs on page 12 of the specification (the second paragraph wraps around to the top of page 13) with the following two consecutive paragraphs:

C<sup>5</sup> The comparison of sequences and determination of percent identity and similarity between two sequences can be accomplished using a mathematical algorithm. (*Computational Molecular Biology*, Lesk, A.M., ed., Oxford University Press, New York, 1988; *Biocomputing: Informatics and Genome Projects*, Smith, D.W., ed., Academic Press, New York, 1993; *Computer Analysis of Sequence Data, Part 1*, Griffin, A.M., and Griffin, H.G., eds., Humana Press, New